

Fig. 1A

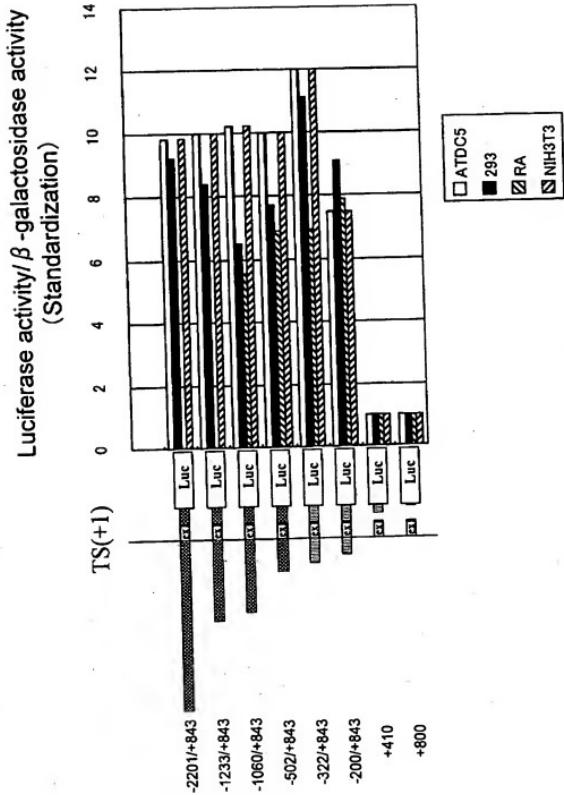


Fig.1B

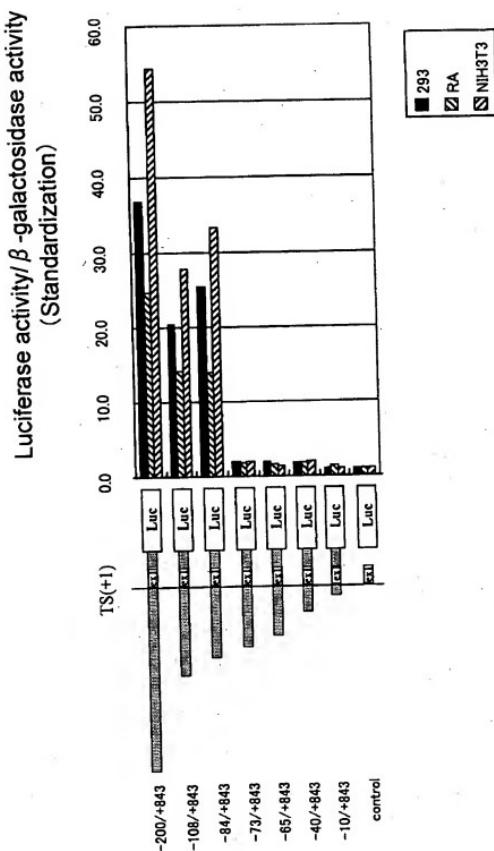
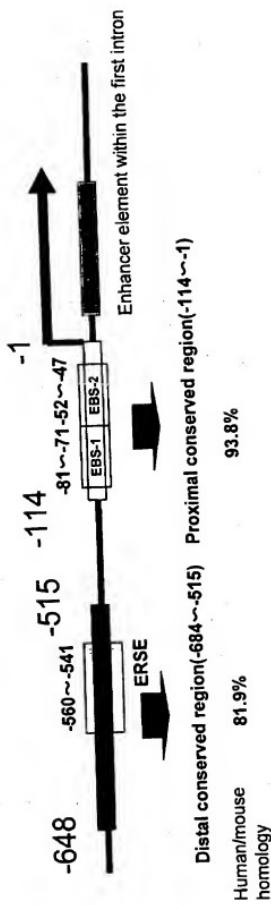


Fig. 1C



96.9% / 97bp

-100

<b>HUMAN</b> CCCACccggcccccaccc ACTGccggcccccaccc	<b>MOUSE</b> GAAATTccgggttccggat GAATTCGGGTTCCGCA	<b>HUMAN</b> CTTACCCCC CTCTACCCCC	<b>MOUSE</b> GCTGAGGG GCTGAGGG
<b>HUMAN</b> -100 SBS-1 EBS-1 ABS	<b>MOUSE</b> -100 SBS-1 EBS-2 SBS-1		
<i>NFI / CTF</i>			

Fig. 1D

Transcription initiation site (+)

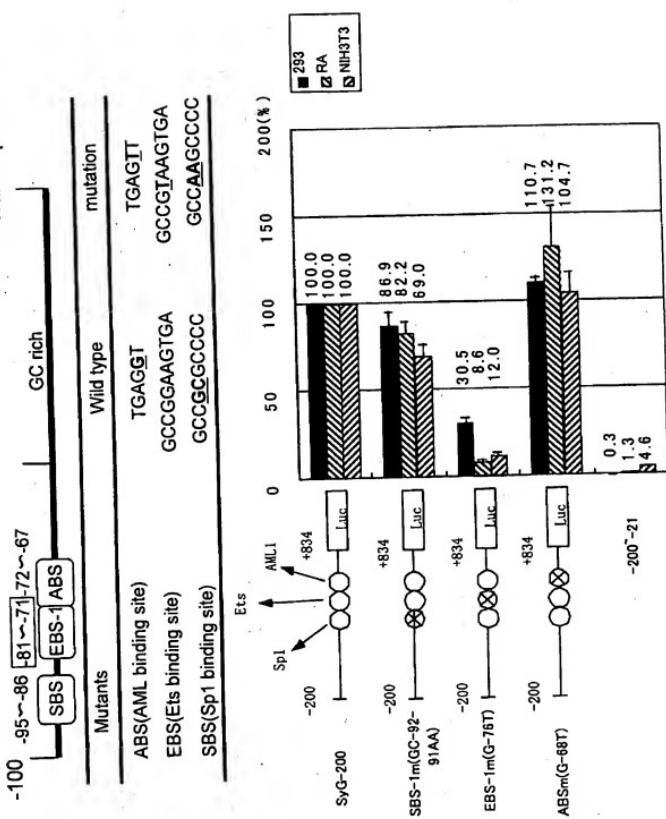


Fig.2A

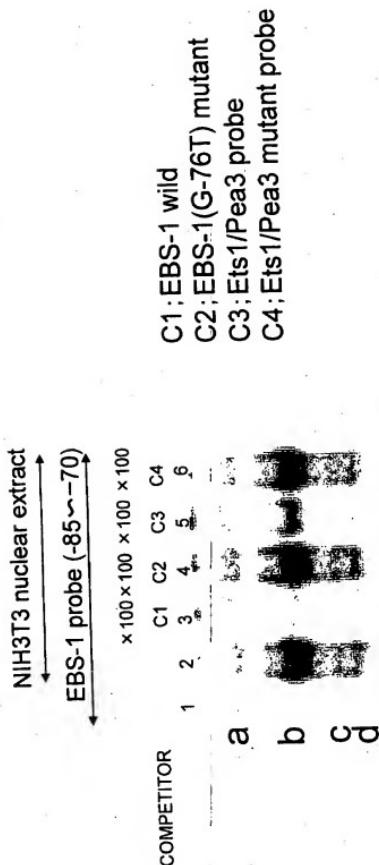


Fig.2B

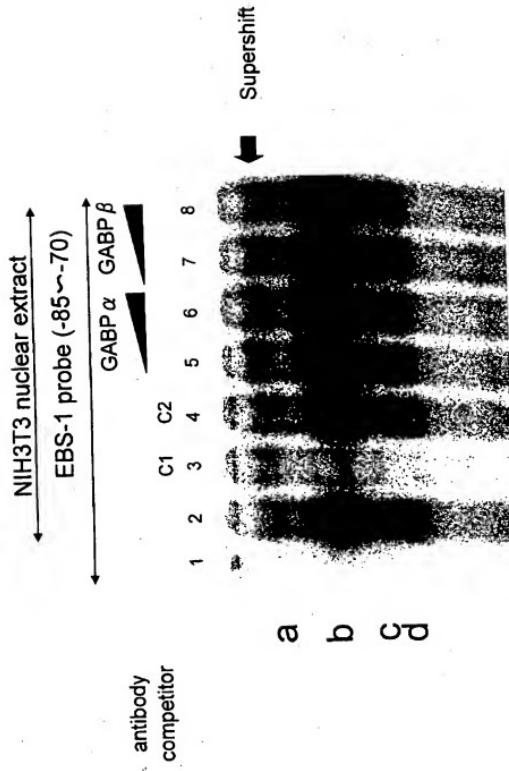


Fig. 2C

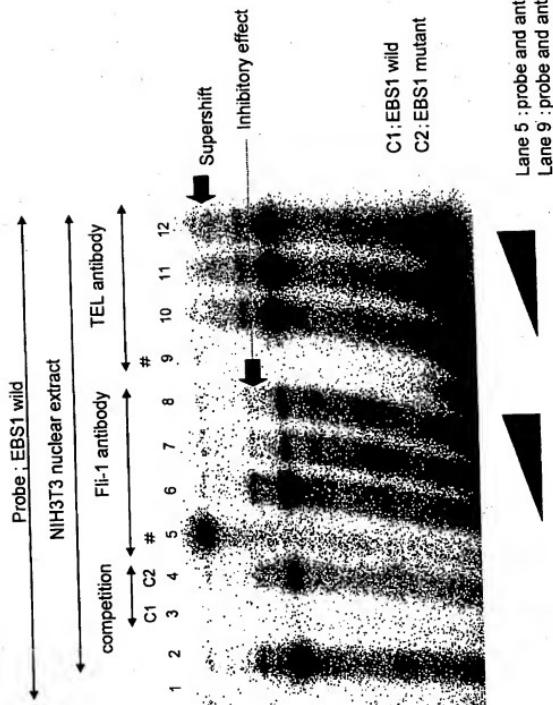


Fig.2D

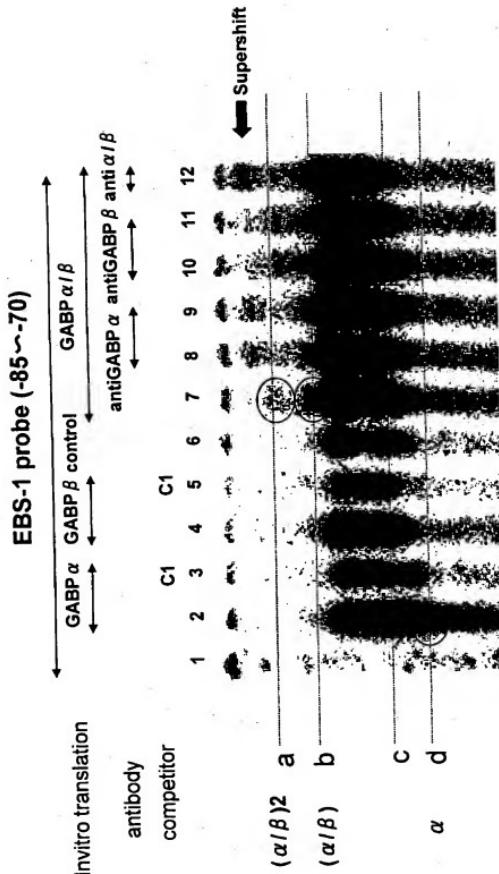


Fig.3A

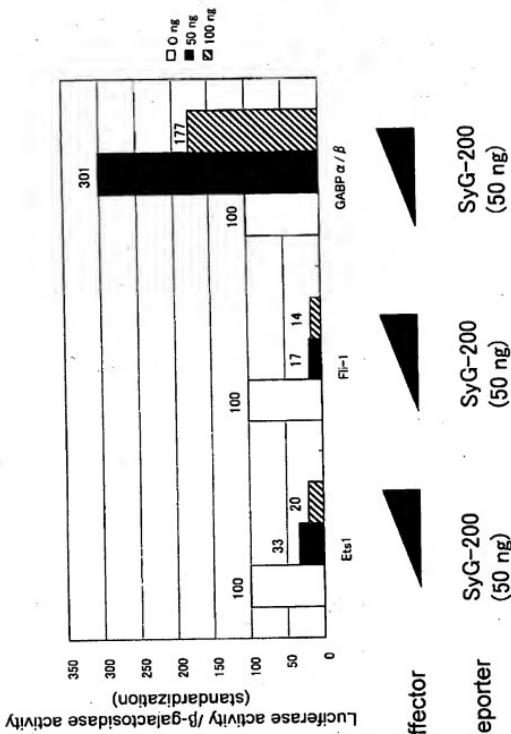


Fig.3B

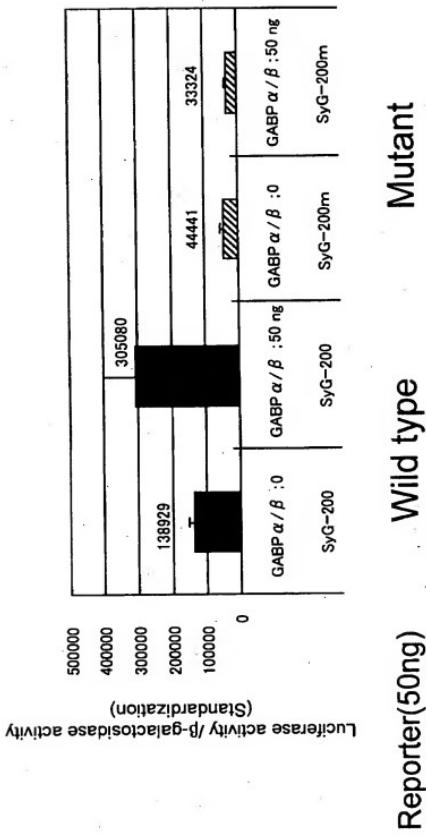


Fig.3C

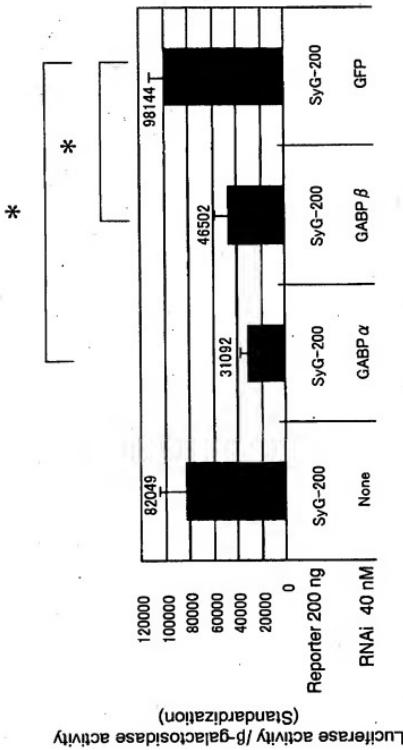


Fig. 4A

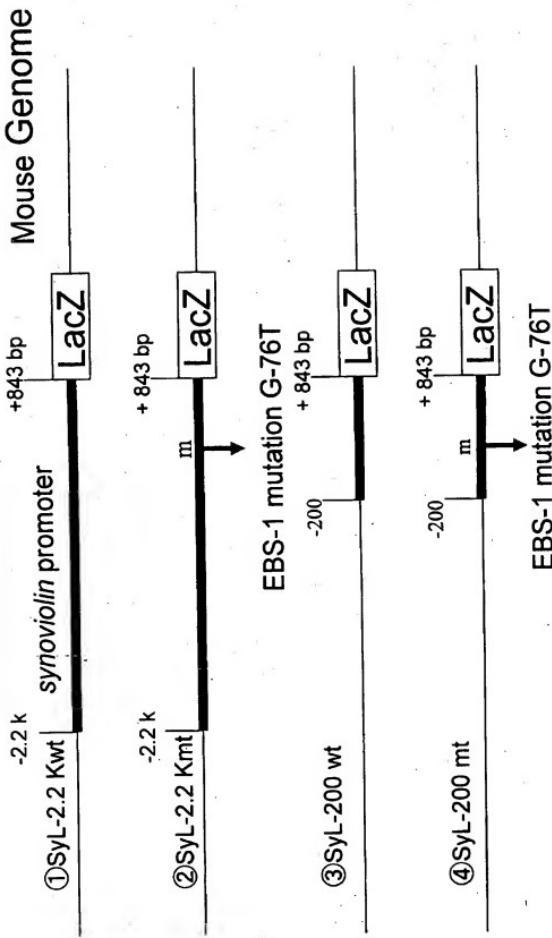


Fig.4B

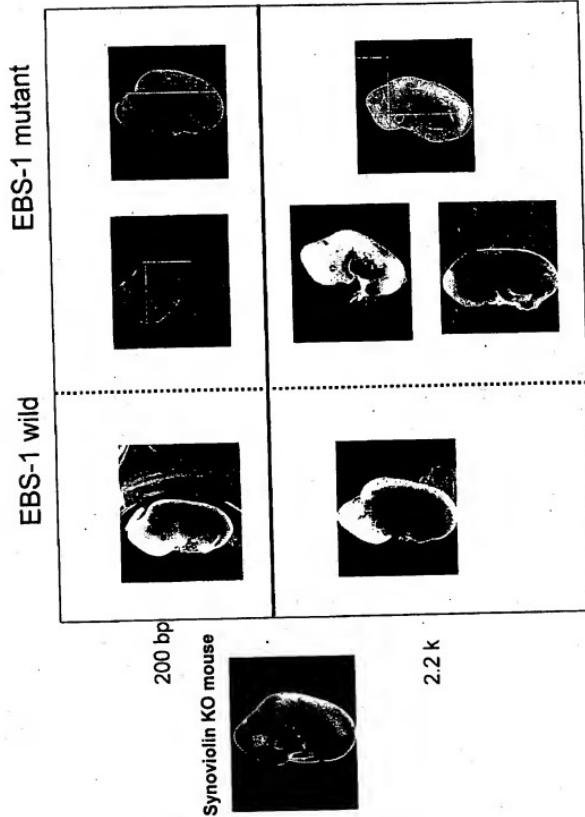


Fig.4C

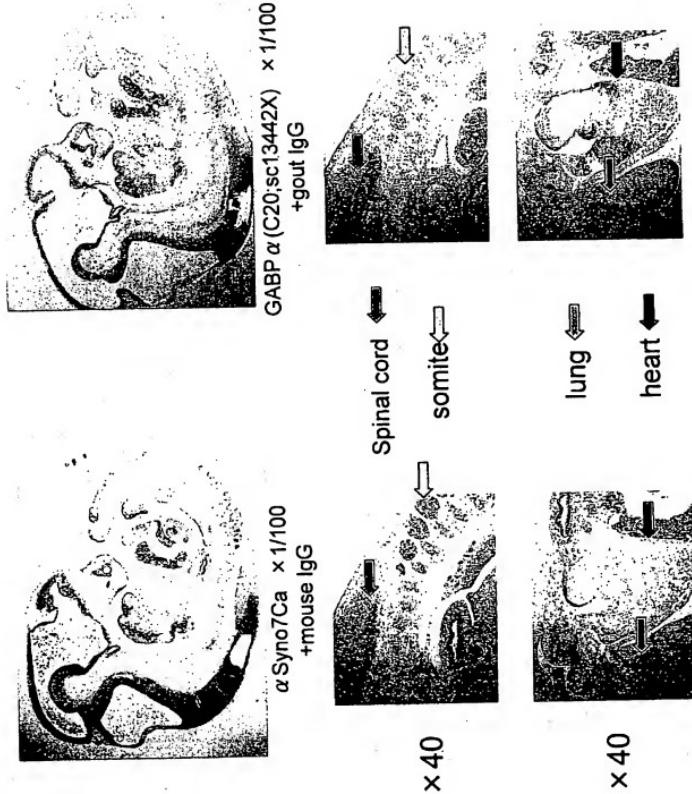


Fig.5

Sequence name	Sequence
EBS-1 Decoy	5'-GCG CCG CGG GAA GTG AGG TG-3' 3'-CGC GGC GGC CTT CAC TCC AG-5'
Scrambled (Negative control)	5'-TTG CCG TAC CCT ACT TAG CC-3' 3'-AAC GGC ATG GGA TGA ATC GG-5'

Fig.6

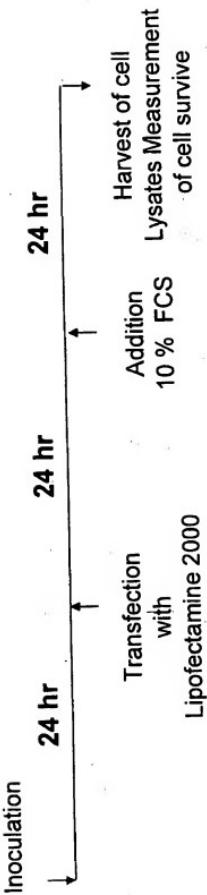


Fig. 7A

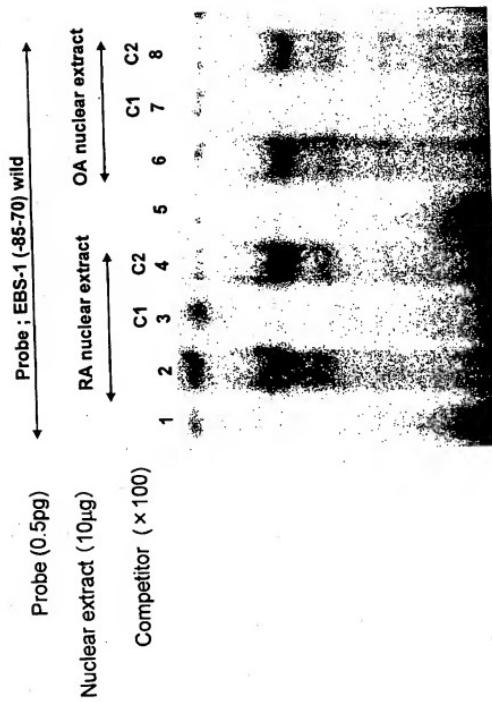


Fig.7B

Cell proliferation activity

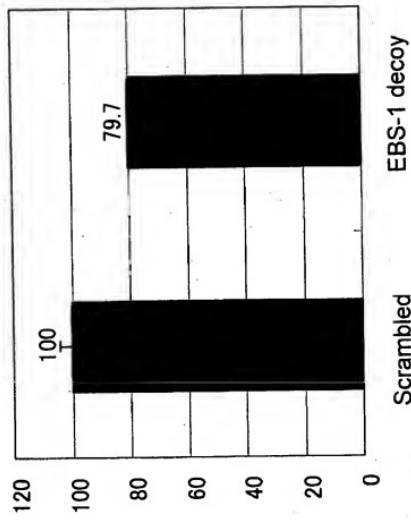


Fig.7C

### Western blotting

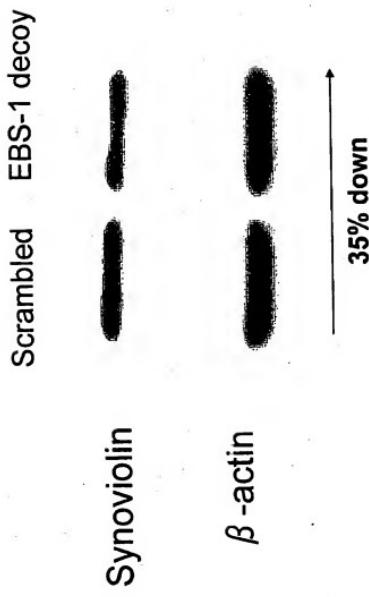


Fig. 8A

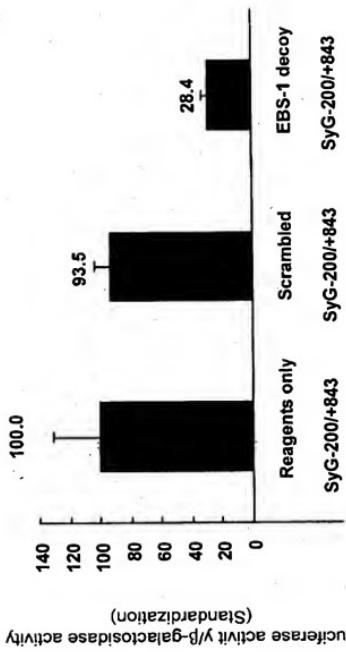


Fig. 8B

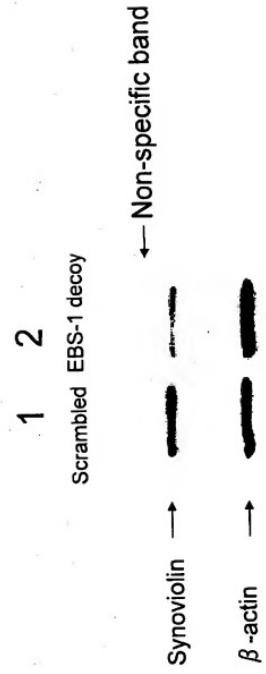


Fig.8C

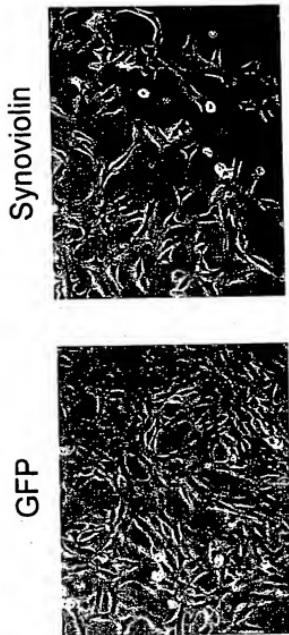


Fig.9A

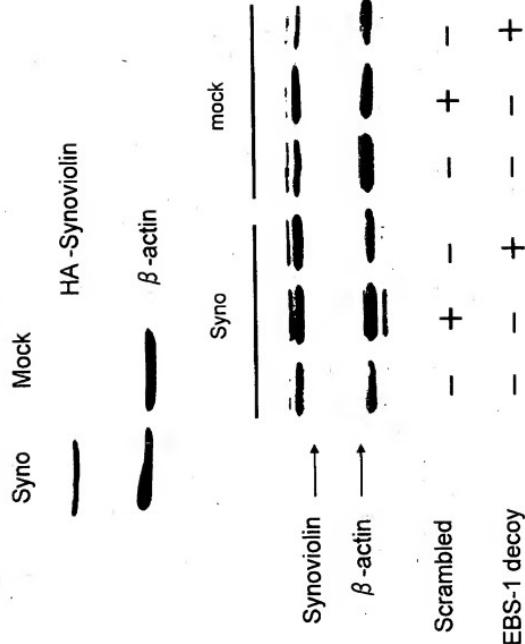


Fig.9B

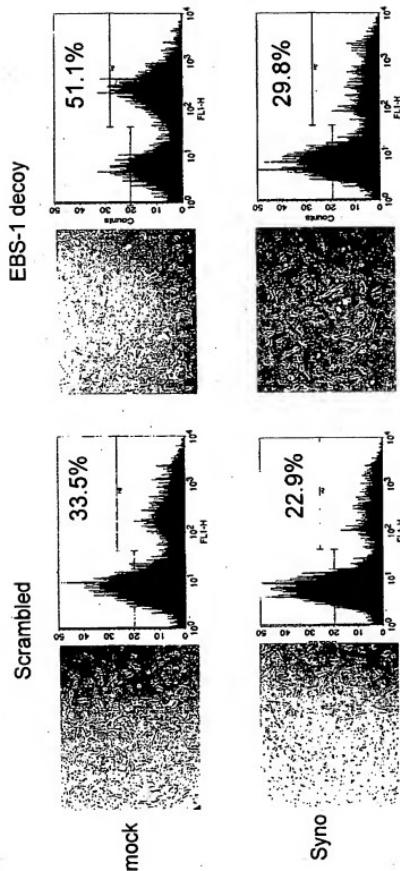


Fig.10

